This Raw Listing contains the General Information Section and up to first 5 pages.

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1    <110> APPLICANT: Brian Seed
2          Tara Pouyani
3    <120> TITLE OF INVENTION: P-SELECTIN LIGANDS AND RELATED MOLECULES
4          AND METHODS
5    <130> FILE REFERENCE: 00786/284002
6    <140> CURRENT APPLICATION NUMBER: US/08/756,018A
7    <141> CURRENT FILING DATE: 1996-11-25
8    <150> EARLIER APPLICATION NUMBER: 60/000,213
9    <151> EARLIER FILING DATE: 1995-06-14
10   <150> EARLIER APPLICATION NUMBER: 08/661,960
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111        tgcccccatc ccgggatgag ctgaccaaga accaggtcag cctgacctgc ctggtcaaag   2040
112        gcttctatcc cagcgacatc gccgtggagt gggagagcaa tgggcagccg gagaacaact   2100
113        acaagaccac gcctcccgtg ctggactccg acggctcctt cttcctctac agcaagctca   2160
114        ccgtggacaa gagcaggtgg cagcagggga cgtcttctc atgctccgtg atgcatgagg   2220
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125                        20                  25                  30
126        Ala Glu Val Lys Lys Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala
127                     35                  40                  45
128        Ser Gly Gly Thr Phe Ser Ser Tyr Ala Ile Ser Trp Val Arg Gln Ala
129                 50                  55                  60
130        Pro Gly Gln Gly Leu Glu Trp Met Gly Gly Ile Ile Pro Ile Phe Gly
131         65                  70                  75                  80
132        Thr Ala Asn Tyr Ala Gln Lys Phe Gln Gly Arg Val Thr Ile Thr Ala
133                         85                  90                  95
134        Asp Glu Ser Thr Ala Arg Asp Asn Gly Ala Tyr Cys Ser Gly Gly Ser
135                        100                 105                 110
136        Cys Tyr Ser Gly Trp Phe Asp Pro Trp Gly Gln Gly Thr Leu Val Thr
137                     115                 120                 125
138        Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro
139                 130                 135                 140
140        Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val
141         145                 150                 155                 160
142        Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala
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150          Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
151          225                    230                    235                    240
152          Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
153                        245                    250                    255
154          Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
155                    260                    265                    270
156          Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
157                    275                    280                    285
158          Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
159              290                    295                    300
160          His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
161          305                    310                    315                    320
162          Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
163                    325                    330                    335
164          Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
165                    340                    345                    350
166          Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
167                    355                    360                    365
168          Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
169              370                    375                    380
170          Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
171          385                    390                    395                    400
172          Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
173                    405                    410                    415
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193       tcggatcccg agggtgagta ctaagcttca gcgctcctgc ctggacgcat cccggctatg    660
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225                    35                  40                  45
226        Ala Phe Arg Asn Glu Glu Tyr Asn Lys Ser Val Gln Glu Ile Gln Ala
227                50                  55                  60
228        Thr Phe Phe Tyr Phe Thr Pro Asn Lys Thr Glu Asp Thr Ile Phe Leu
229        65                  70                  75                  80
230        Arg Glu Tyr Gln Thr Arg Gln Asp Gln Cys Ile Tyr Asn Thr Thr Tyr
231                        85                  90                  95
232        Leu Asn Val Gln Arg Glu Asn Gly Thr Ile Ser Arg Tyr Val Gly Gly
233                        100                 105                 110
234        Gln Glu His Phe Ala His Leu Leu Ile Leu Arg Asp Thr Lys Thr Tyr
235                    115                 120                 125
236        Met Leu Ala Phe Asp Val Asn Asp Glu Lys Asn Trp Gly Leu Ser Val
237                130                 135                 140
238        Tyr Ala Asp Lys Pro Glu Thr Thr Lys Glu Gln Leu Gly Glu Phe Tyr
239        145                 150                 155                 160
240        Glu Ala Leu Asp Cys Leu Arg Ile Pro Lys Ser Asp Val Val Tyr Thr
241                        165                 170                 175
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